

Assessing understanding of complex learning outcomes and real-world skills using an authentic software tool: a study from Biomedical Sciences

Practitioner Research
In Higher Education
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University of Cumbria
Vol 8 (1) pages 101-112

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Abstract

We describe a study conducted during 2009-12 into innovative assessment practice, evaluating an assessed coursework task on a final year Medical Genetics module for Biomedical Science undergraduates. An authentic e-assessment coursework task was developed, integrating objectively marked online questions with an online DNA sequence analysis tool (BLAST), routinely used by NHS and research professionals. The aim was to combine the assessment of understanding of complex module learning outcomes with real-world authentic skills highly valued in the work place. This approach challenges the oft-heard accusation that online computer-marked tests can lack validity and authenticity in higher education. The study demonstrates the content and construct validity of this form of e-assessment, showing that careful question design, allied with integration with the real life BLAST tool, enables instructors to assess complex higher order understanding, and requires students to demonstrate skills relevant for the work place. A study of three years of test results and measures of internal consistency data also show the reliability of this assessment. In addition, the results of surveys of student opinion and positive feedback from student module feedback questionnaires suggest that it is effective in terms of face validity.

Keywords

Authentic assessment; technology enhanced assessment; assessing deeper learning.

Background and Rationale

For some time in higher education, students have been calling for innovative assessments which focus on understanding and application of knowledge instead of memorised techniques (National Student Forum, 2009; National Union of Students, 2009), and which require learners to engage in appropriate learning tasks (Boud et al., 2010). This is especially true of final year students, whose learning outcomes will tend to concentrate on higher order cognitive skills (Bloom, 1956).

However, when there are large numbers of students on a module cohort, it can be a challenge to design assessments which are reliable, valid and practical (Brown and Knight, 1994). In addition, whilst it is well known that multiple choice questions can easily and flexibly be delivered to large numbers of students, and these can be automatically objectively marked and graded reliably, with instant results and feedback given to students (Bull and McKenna, 2004; Crisp, 2007), there is some debate in the literature as to whether these objectively marked questions are only limited to testing lower order skills, and it has been argued that these question types need to be used more imaginatively to engage students in the assessment and learning process (Nicol, 2007; Nicol and Macfarlane-Dick, 2006; Gibbs and Simpson, 2004).

Historically, computer assisted assessment in the biosciences has tended to comprise paper-based multiple choice questions assessing student knowledge and understanding at a more superficial level. However, numerous recent innovative projects and high-profile support for e-learning from JISC (2007; 2010) have demonstrated a wide range of benefits offered by e-assessment, many of

Citation

Dermo, J., Boyne, J. (2014) Assessing understanding of complex learning outcomes and real-world skills using an authentic software tool: a study from biomedical sciences, *Practitioner Research in Higher Education Journal*, 8(1), January, 101-112.

which are directly appropriate to this study and include: greater variety and authenticity in the design of assessments, capture of wider skills and attributes not easily assessed by other means, efficient submission, marking, moderation and data storage processes (Bryan and Clegg, 2006; Jordan, 2013).

In this case study we were interested in developing a problem-based assessment that replicates the kind of investigatory data analysis a medical geneticist might undertake when diagnosing a patient and thus represented a 'hands-on' type of learning exercise (Sivan et al., 2000). Genetic disorders are routinely diagnosed by DNA sequence analysis. Raw DNA sequence is essentially a series of letters (A, T, C and G – the genetic 'code') that are generally analysed using on-line software called Basic Local Alignment Search Tool (BLAST). The BLAST software is operated by The National Centre for Biotechnology Information (NCBI), which advances science and health by providing free access to biomedical and genomic information. The way most people use BLAST is to input a DNA sequence as a query against all of the public sequence databases, pasting the sequence into the textbox on one of the BLAST Web pages. This sends the query over the Internet, the search is performed on the NCBI databases and servers, and the results are posted back to the person's browser in the chosen display format, usually within sixty seconds of submitting the query (Madden, 2002). Experience with this tool represents an applied skill that would directly benefit graduates who chose to enter a scientific career involving genetic analysis (a discipline that expands year-on-year across the research, diagnostic and pharmaceutical sectors) and also provides an excellent interactive tool around which to frame a problem-based assessment that tested the students' knowledge and understanding of the modules learning outcomes.

The aim was to create a coursework test for this module which met the following requirements: first, it must be practical, able to be delivered to large numbers of students, marked automatically and administered by a small module team; second, this must be a valid test, in terms both of content validity, with the assessment constructively aligned with the learning outcomes of a year three module (Biggs, 1999), and in terms of face validity, since it is important for the students to consider the assessment to be a 'good' test (Dermo, 2009). In addition, the test must be secure and reliable. The major logistical challenges to be addressed during development of the assessment were ensuring that students who had access to the Internet (BLAST runs via browser and requires an active Internet connection) did not retrieve material that would jeopardise test security and invalidate the summative assessment.

The research questions which this study aimed to answer were:

- Was it possible to devise a test which could be delivered securely and administered economically to approximately 150 students, and marked automatically?
- Was this test a valid and authentic assessment of the learning objectives for the level 6 (i.e. final year undergraduate) module, challenging the students at an appropriate level?
- Was this test a reliable measure of achievement?
- Was the test viewed by the students as a valid assessment at their level?

Assessment Tools and Processes

The University of Bradford operates a dedicated e-assessment suite, designed to run high-stakes assessment via a thin-client based server array (Richardson et al., 1998; Dermo, 2011). One of the advantages of this system is that terminals can be modified so that web browsers lack address toolbars and open directly at the NCBI homepage, thus confining student access to the BLAST software during the summative assessment. Moreover, these terminals are linked to a secure server that enables students to resume assessments on any of the other terminals in the suite should a terminal fail and also serves to provide high-stakes encryption of assessment data.

Questionmark Perception (QMP) is the University of Bradford's chosen tool for summative high-stakes e-assessment. It is widely used for formative and summative online assessments, supports varied question types, can deliver generic pre-prepared feedback on a question level or by topic and is an established tool for assessment and feedback in the biosciences (Olson and McDonald, 2004). Many of these question types are closed questions, which can be automatically marked by the system enabling rapid turnaround and same-day release of marks to students.

In order to introduce students to the e-assessment and familiarise them with the BLAST tool, formative exercises related to the assessment were made available via the University of Bradford's virtual learning environment (Blackboard). This enabled students to practise both the style of questions that would appear in the summative assessment and gain some self-directed experience in using BLAST throughout the module. Formative micro-assessments such as this have a proven track record in Bioscience degrees (Thin, 2006). This continuous assessment process was supported by a dedicated formative workshop in the e-assessment suite where students had the opportunity to attempt formative questions and interface with the BLAST tool in a manner consistent with the summative exam, thus gaining some insight into how their learning was developing (Ramsden, 1999). This is good practice as laid down by the University's policy for computer assisted assessment.

The summative e-assessment comprises forty closed questions delivered online: various question types were used, including drag and drop, fill in the blanks, hotspot, matching, multiple choice, multiple response, numeric, pull-down list, ranking and select-a-blank. These questions relate to 'genetics case studies' that present the student with DNA sequence and task them to analyse this information using the BLAST tool and use the data to answer the questions. This format, therefore, establishes a process whereby the learner is required to apply knowledge and expertise when interpreting the results of the BLAST search tool combined with knowledge and understanding of the module learning outcomes. An example of a case study and related question can be seen in Figure 1.

Questionmark

A 'nucleotide BLAST' search was performed using the following sequence:
TGTGTCTGAACTGGCTGCTGAATGTTTATG

The results of the BLAST search are shown below:

Alignments

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Homo sapiens dystrophin (DMD), transcript variant Dp140c, mRNA
 Sequence ID: refNM_004020.3 Length: 7080 Number of Matches: 1

Range 1: 3191 to 3220 GenBank Graphics

Score	Expect	Identities	Gaps	Strand
60.0 bits(30)	5e-07	30/30(100%)	0/30(0%)	Plus/Plus

Query 1 TGTGTCTGAACTGGCTGCTGAATGTTTATG 30
 |||
 Sbjct 3191 TGTGTCTGAACTGGCTGCTGAATGTTTATG 3220

Download ▾ GenBank Graphics

Homo sapiens dystrophin (DMD), transcript variant Dp140bc, mRNA
 Sequence ID: refNM_004023.2 Length: 7048 Number of Matches: 1

Range 1: 3191 to 3220 GenBank Graphics

Score	Expect	Identities	Gaps	Strand
60.0 bits(30)	5e-07	30/30(100%)	0/30(0%)	Plus/Plus

Query 1 TGTGTCTGAACTGGCTGCTGAATGTTTATG 30
 |||
 Sbjct 3191 TGTGTCTGAACTGGCTGCTGAATGTTTATG 3220

Which of the following mutations is most commonly associated with this gene?

- Missense
- Deletion
- Nonsense
- Chromosomal translocation
- Tri-nucleotide repeat expansion

Submit

Figure 1. Screen capture of the BLAST software data and associated multiple choice question taken from a previous iteration of the e-assessment. Note the requirement to be able both to interpret the BLAST data and possess module specific knowledge relating to that data in order to correctly answer the question.

As it was originally envisaged and designed, the online test itself consists of two simultaneous browser sessions. In one browser window, the students have access to the actual BLAST tool. In the second browser, the students are delivered an online test in which they are given a DNA sequence, which they copy and paste into the BLAST tool for analysis. Students are then shown a number of questions about the genetic data they have just inputted into BLAST. The students have to read and understand the data sent back to them by BLAST and answer the questions. In this way the students have to be able to use the BLAST tool appropriately, and understand and interpret the data sent back to them. This replicates closely how the BLAST tool is used in the work place. Although in the real world they would be using the data to inform decisions, not to answer MCQs, the students are applying their knowledge of the subject to be able to answer the questions correctly and this can certainly be considered to be an authentic assessment task (Boud, 2000; Boud and Falchikov, 2006).

The assessment has been administered over three academic years and item analysis has routinely been run on item performance. After use, items are kept secure in a virtual item bank and new questions are added to the bank each year. The test was deliberately designed so that new questions could be quickly and easily developed, based on the learning outcomes. It is anticipated that soon a complete and comprehensive bank of secure items will be established for use in future years. Of course, quality assurance checks are carried out in conjunction with external examiners, as

with any high stakes assessment items, according to the UK Quality Code for Higher Education (QAA, 2011).

Methods

For this case study, test result statistics and student questionnaire data were gathered and analysed in order to answer the research questions above. As is the case with many educational studies, a mixed methods approach is appropriate, combining qualitative and quantitative approaches to data collection and analysis (Pring, 2004).

Quantitative Data Analysis

Three years of coursework scores were collected from 2010 to 2013. A post hoc analysis of these data was carried out to investigate whether there was an acceptable distribution of scores and mean scores which would indicate that the assessment has been sufficiently challenging for the students, in line with normal procedures for quality assurance. Objectively marked questions have been criticised for not assessing at a high enough level for higher education: if these tests can show a high enough mean score and can approach normal distribution, then these criticisms can be challenged. In addition, the reliability of the assessment was measured with a test of internal consistency (Cronbach's Alpha).

Qualitative Data Analysis

A specific student questionnaire was delivered to a student cohort (in year 2 of the study). This consisted of closed Likert scale opinion questions, using the scale strongly agree / agree / neutral / disagree / strongly disagree, as well as open-ended questions to elicit comments on positive aspects as well as constructive criticism. The questions used in the student questionnaire were selected based on the issues arising from the literature, as outlined above, and can be seen in Figure 2. below:

1. Learning to use the BLAST software to analyse DNA sequences is well suited to the e-assessment facilities at UoB
2. The extended MCQs tested my understanding (as opposed to simply recollection) of the module material
3. This e-assessment assesses things that would not have been possible on a paper-based test
4. The use of BLAST and this e-assessment are simply a gimmick and do not benefit my learning.
5. Do you have any positive comments about this assessment?
6. Have you any suggestions how this assessment can be improved?

Figure 2. Questions used for student qualitative data questionnaire.

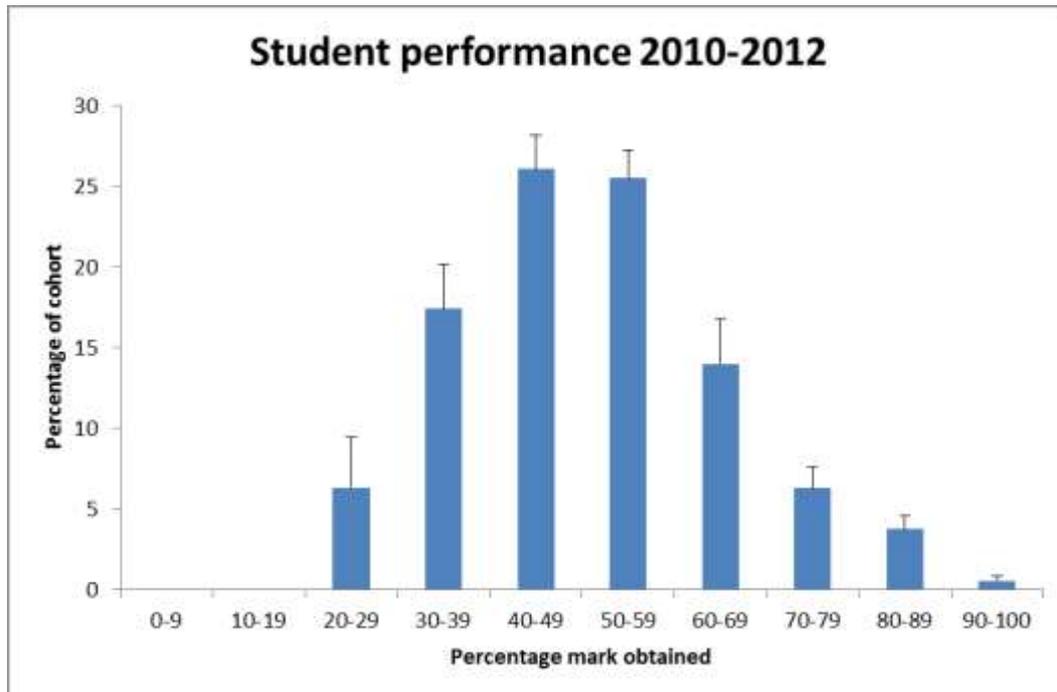
In addition, the researchers were able to search through data from the university module evaluation questionnaires completed by students over the course of the three years. This generic questionnaire does not specifically address the BLAST assessment, but there is space for additional student comments, where we might expect to find references to the assessment on the module.

Results and Findings

Quantitative Data Analysis

The first iteration of the summative e-assessment ran in November 2010 and has run for three academic cycles. In order to compare student performance in the e-assessment across all three iterations, the distribution of marks for each cohort was calculated in relation to the percentage of students from each year and mean values plus standard error of the mean derived. These data can be seen in Figure 3a., which demonstrates a normal-like distribution of marks. Similarly, the mean

mark obtained in the e-assessment over three academic years was calculated alongside standard deviation and standard error of the mean (Figure 3b.).



	2010	2011	2012
Mean	48.78	50.85	52.08
SD	13.02	13.91	14.12
SEM	1.14	1.14	1.06

Figure 3. Histogram showing distribution of percentage coursework scores 2010-2013 and a table showing mean percentage scores by year, with standard deviation and standard error of measurement.

The distribution of coursework scores for each of the three different cohorts of students in this study also approximate normal distribution, with mean scores ranging from 48.8% to 52.1% (see Figure 4.).

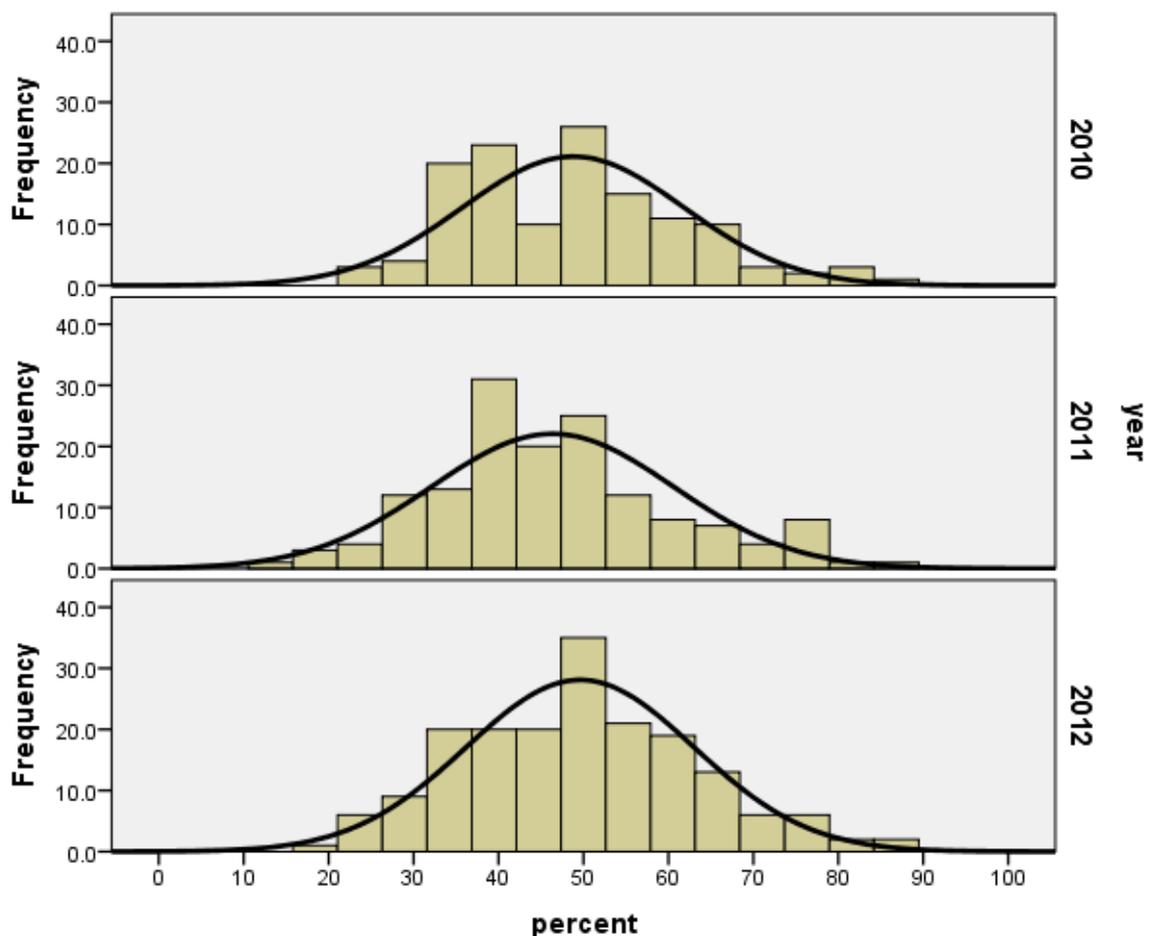


Figure 4. Histograms showing the performance of each year’s student cohort, indicating a normal-like distribution of scores and consistent performance from year to year.

In terms of inter-item reliability, the test performed well from year to year, with a Cronbach’s alpha value between 0.7 and 0.8 for each administration of the test (see figure 5.). This would generally be considered as a good rating as an internal consistency estimate of reliability of the test scores (Kline, 1999).

Test reliability: Cronbach's Alpha	
2010-11	0.77
2011-2	0.78
2012-3	0.70

Figure 5. Assessment reliability 2010-2013.

Qualitative Data Analysis

A student questionnaire was distributed to the 2011-2 student cohort, with a response rate of over 75 per cent ($n = 115/150 = 76.67\%$). Collated student responses, (as outlined in Figure 6.) reveal that a large majority of respondents (83.5%) agreed or strongly agreed that the BLAST test was definitely well suited to the facilities. Almost three-quarters (72.2%) or students stated that they believed that the questions assessed understanding rather than factual recall, and approximately two thirds (64.4%) were of the opinion that e-assessment can enable activities which would not be possible on paper. A similar number (70.4%) believe that this assessment was more than a gimmick and does benefit learning.

	Strongly Agree	Agree	Neutral	Disagree	Strongly Disagree
Learning to use the BLAST software to analyse DNA sequences is well suited to the e-assessment facilities at the UoB.	29.6	53.9	10.4	2.6	3.5
The extended MCQs tested my understanding (as opposed to simply recollection) of the module material.	25.2	47	17.4	5.2	5.2
This e-assessment assesses things that would not be possible on a paper-based test.	18.3	46.1	19.1	12.2	4.3
The use of BLAST and this e-assessment are simply a 'gimmick' and do not benefit my learning.	3.5	7	19.1	39.1	31.3

Figure 6: Percentage responses to Likert scale opinion questions on student questionnaire.

In addition, respondents were able to write comments in response to the open-ended prompt 'Do you have any positive comments?' These free responses were analysed and collated and the most frequent themes were identified.

Students mentioned that the test was a good assessment of understanding and application of knowledge, which concurs with the second question on the questionnaire. It was also stated that the test was well organized, clear and easy to use, and it was recognized that marking and feedback are quicker. Students found the process helpful for their revision and study and valued the BLAST tool as useful to them. They also appreciated seeing a new approach to assessment and some liked the fact that the task was challenging for them. Figure 7. contains some examples of the positive comments raised by students to the open-ended questions.

Some comments from students:

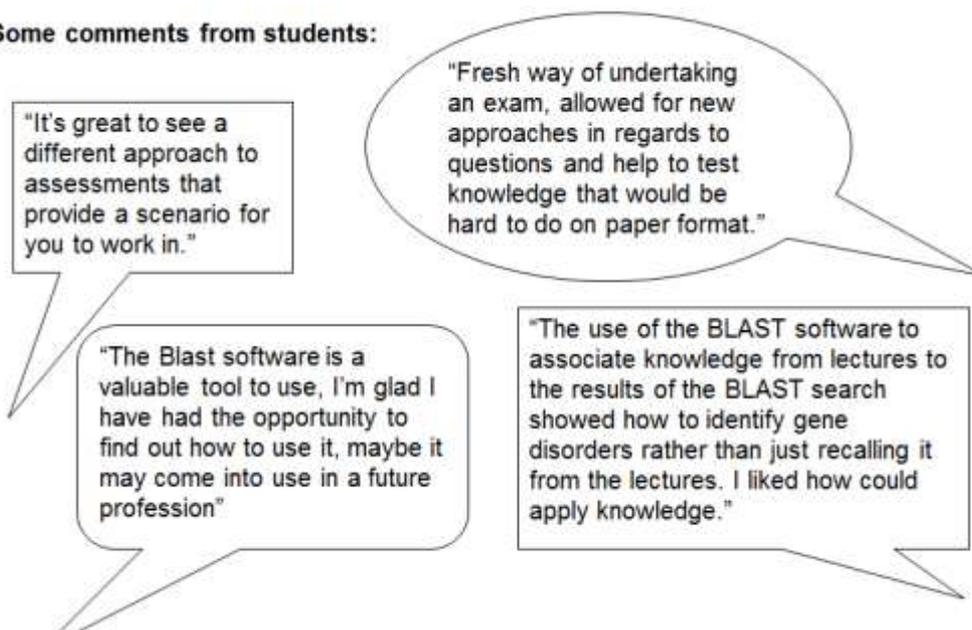


Figure 7: Some quotes from students, typifying some of the positive themes emerging from open student responses on the questionnaire.

In addition, respondents were asked to offer constructive criticism. The most frequent suggestions and criticisms were related to the fact that some students found the test too difficult, or that it covered too much content. There were also some comments related to technical issues related to the BLAST tool or IT issues in general, especially concerning slow response times, waiting and the pressure and stress which this caused.

The researchers also examined the completed general module evaluation feedback questionnaires for the Medical Genetics module over three years. Only a few such comments were identified, and these typically gave positive reflections on the experience, reiterating the positive themes emerging from the questionnaire responses. Figure 8 shows typical comments which were found.

'The coursework elements were well received. Using the blast software was something novel and never experienced before in a degree capacity.'

'The individual assessment ran quite well on the computers and it also allowed for quick feedback to be gained in terms of results.'

'20% computer based examination is a lot better than MCQ based exam as it taught me how to use BLAST well. I like the fact that we are examined in different area this year.'

'I enjoyed most of the lectures, particularly enjoyed learning about diseases. The first assessment was difficult but it really pushed me to go through the lecture slides.'

Figure 8. Extracts from module feedback questionnaires.

Discussion

Student performance and qualitative feedback broadly demonstrate that the e-assessment has met expectations with regard to providing a challenging and innovative assessment and providing students with discipline-relevant skills.

In terms of the research questions specified earlier, we can conclude that it is possible to devise a test which can be delivered securely and administered economically to large groups of students. Using an authentic online research tool alongside online questions, this test provides a reliable, valid and authentic assessment of the learning objectives for the module, offering a challenging assessment of their skills. In addition, qualitative data supports the idea that the test was certainly viewed by the students as a valid assessment.

However, these data also highlight several areas where improvement and modification can be made, especially with regard to learner support and risk management.

Arguably, the most innovative aspect of the e-assessment presented in this article is the interface between data obtained using the BLAST software and online questions, which requires students to interpret BLAST data and link it to module-specific learning outcomes. A greater emphasis on the problem-solving aspect of questions and relying less on the learner's ability to recall key terms and phrases from lectures is something we wish to implement in future versions of the e-assessment. One route for this is to train students to use the BLAST software beyond basic DNA sequence analysis. This would enable far more complex case studies to be written that include problems which

require participants to navigate through the BLAST software in a more involved manner in order to obtain the data relevant to the question being answered. Such questions are inherently more interactive in nature and begin to approach 'real-world' scenarios, two elements that others have shown to both motivate students and encourage strong independent learning (Mustoe and Croft, 1999). We are also interested in the possibility of running formative team-based learning (TBL) sessions around this assessment. Such collaborative learning environments would be well suited to the formative session that aims to develop student proficiency with the BLAST software. There is strong evidence that demonstrates such TBL sessions foster active learning and improves critical thinking amongst students (Allen and Tanner, 2005; Herreid, 2013), two goals that are consistent with the overarching aims of the e-assessment.

The mean mark for the BLAST e-assessment for the three cohorts is 50.57%, which demonstrates that students find the assessment somewhat challenging. While the Cronbach Alpha coefficients suggest that internal consistency of the questions is sound, it is important to reflect upon the ratio of different question types used throughout the assessment and adjust these so that the test remains challenging, but enables a normal distribution of marks. As can be seen in Figure 3a., the data gathered on student performance over three years approaches a normal distribution, but there is an overrepresentation of candidates in the 40-49% bracket of marks. Currently, of the forty questions that comprise the summative assessment, fifteen of these are multiple response questions (MRQs), where students need to select more than one option to gain full marks. Such question types are valuable when attempting to generate more authentic objectively-marked assessments, however, there is also a risk that the complex nature of MRQs may lead to an inability to discriminate between weaker and stronger students and thus negatively impact on test quality (McAlpine and Hesketh, 2003). On reflection, we feel that MRQs may be slightly overrepresented in the e-assessment and intend to reduce the number of these question types from fifteen down to ten in the 2013 iteration of the test.

As discussed earlier, the University of Bradford's high-stakes assessment suite utilises a thin-client based server array that enables restricted access to websites. One unfortunate caveat of using the BLAST software is that the NCBI website also contains pages with significant information about the structure, function and medical relevance of genes. This means that allowing students access to NCBI website to use the BLAST tool also permits them to see material that (if accessed) would invalidate the summative closed-book assessment. Currently, these risks are mitigated by invigilation by four academics, one for every 25 students, but we are keen to explore other options. Crucially, the work-relevant aspects of interpreting data derived from the BLAST software needs to remain *in situ* as losing this would undermine the remit of the e-assessment and render it unfit for purpose.

A second risk management issue relates to the response time following a request being sent to the BLAST server. Student feedback provided in the questionnaire raised concerns over this process being both lengthy and unpredictable. The length of time it takes BLAST to return data can vary between a few seconds to a few minutes, depending on the complexity of the requested search and the time of day. Slower response times reflect peak-usage and this tends to coincide with daylight hours in the U.S. Obviously, it is imperative that all students have an equivalent assessment experience and therefore these concerns need to be addressed.

One possible solution to both of the issues discussed above would be to present students with 'screen capture' images taken from the BLAST software during the summative assessment, rather than requiring the student to access the BLAST data online. Such an approach would do little to reduce the validity and authenticity of the assessment, as the key skills associated with the BLAST software are in data analysis, rather than the actual entry of query sequence, the latter simply requiring student to 'copy and paste' a string of letters into the BLAST search field. This change

would serve to remove any variation in assessment experience caused by varying BLAST search times and remove the possibility of candidates accessing inappropriate materials via the Internet during the exam. It would also reduce the amount of software running on the thin-client server from two (online test and web browser with BLAST) to one, which is likely to improve the stability of the e-assessment. Were this change to be implemented, we would likely improve on the formative BLAST session, perhaps by switching to a team-based learning approach, in order to use this time to develop student proficiency with the 'live' BLAST software so that this aspect of the e-assessment is improved, rather than diminished due to the 'offline' nature of the summative exam.

One other area for development for this assessment is to create a mobile learning version of the practice version of this assessment, using rich formative feedback to support the students during the learning process. Data gathered on the Medical Genetics module as part of an HEA-funded Individual Teaching Development project has indicated considerable interest in such a tool among these students, and work is already underway to develop and implement a novel, interactive mobile learning resource that students will be able to access any time, in any surrounding in order to engage with the module learning outcomes.

Conclusion

In conclusion, it has been possible to use the BLAST tool and Questionmark Perception online assessments to devise a test which can be delivered securely and administered economically to large groups of students, and marked automatically. In addition, there is evidence that this test is a valid, authentic and reliable assessment of the learning objectives for final year undergraduates which challenges the students at an appropriate level. The results of surveys of student opinion, and positive feedback from student module feedback questionnaires also suggest that the test is effective in terms of face validity.

The conclusions from this study are of potential interest and relevance to lecturers across a range of disciplines or professional fields: in particular, careful design of online assessment questions, in conjunction with integration with real life authentic online tools, can enable instructors to assess complex higher order understanding in a valid, reliable and practical way, and can require students to demonstrate skills relevant for the work place.

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